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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
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EXAMINER

GOLDBERG, JEANINE ANNE

ART UNIT

PAPER NUMBER

1634

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PAPER

**Please find below and/or attached an Office communication concerning this application or proceeding.**

The time period for reply, if any, is set in the attached communication.

<b>Office Action Summary</b>	<b>Application No.</b> 10/677,701	<b>Applicant(s)</b> LEVENSON ET AL.	
	<b>Examiner</b> JEANINE A. GOLDBERG	<b>Art Unit</b> 1634	

-- The MAILING DATE of this communication appears on the cover sheet with the correspondence address --

### Period for Reply

A SHORTENED STATUTORY PERIOD FOR REPLY IS SET TO EXPIRE 3 MONTH(S) OR THIRTY (30) DAYS, WHICHEVER IS LONGER, FROM THE MAILING DATE OF THIS COMMUNICATION.

- Extensions of time may be available under the provisions of 37 CFR 1.136(a). In no event, however, may a reply be timely filed after SIX (6) MONTHS from the mailing date of this communication.
- If NO period for reply is specified above, the maximum statutory period will apply and will expire SIX (6) MONTHS from the mailing date of this communication.
- Failure to reply within the set or extended period for reply will, by statute, cause the application to become ABANDONED (35 U.S.C. § 133). Any reply received by the Office later than three months after the mailing date of this communication, even if timely filed, may reduce any earned patent term adjustment. See 37 CFR 1.704(b).

### Status

- 1) ☒ Responsive to communication(s) filed on 10 September 2007.
- 2a) ☒ This action is **FINAL**.                      2b) ☐ This action is non-final.
- 3) ☐ Since this application is in condition for allowance except for formal matters, prosecution as to the merits is closed in accordance with the practice under *Ex parte Quayle*, 1935 C.D. 11, 453 O.G. 213.

### Disposition of Claims

- 4) ☒ Claim(s) 1-5, 12-14, 23-28, 31, 33 and 34 is/are pending in the application.
- 4a) Of the above claim(s) \_\_\_\_\_ is/are withdrawn from consideration.
- 5) ☐ Claim(s) \_\_\_\_\_ is/are allowed.
- 6) ☒ Claim(s) 1-5, 12-14, 23-28, 31, 33 and 34 is/are rejected.
- 7) ☐ Claim(s) \_\_\_\_\_ is/are objected to.
- 8) ☐ Claim(s) \_\_\_\_\_ are subject to restriction and/or election requirement.

### Application Papers

- 9) ☐ The specification is objected to by the Examiner.
- 10) ☐ The drawing(s) filed on \_\_\_\_\_ is/are: a) ☐ accepted or b) ☐ objected to by the Examiner.  
Applicant may not request that any objection to the drawing(s) be held in abeyance. See 37 CFR 1.85(a).  
Replacement drawing sheet(s) including the correction is required if the drawing(s) is objected to. See 37 CFR 1.121(d).
- 11) ☐ The oath or declaration is objected to by the Examiner. Note the attached Office Action or form PTO-152.

### Priority under 35 U.S.C. § 119

- 12) ☐ Acknowledgment is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d) or (f).
- a) ☐ All    b) ☐ Some \*    c) ☐ None of:
1. ☐ Certified copies of the priority documents have been received.
  2. ☐ Certified copies of the priority documents have been received in Application No. \_\_\_\_\_.
  3. ☐ Copies of the certified copies of the priority documents have been received in this National Stage application from the International Bureau (PCT Rule 17.2(a)).

\* See the attached detailed Office action for a list of the certified copies not received.

### Attachment(s)

- |  |   |
|--|---|
| 1) <input checked="" type="checkbox"/> Notice of References Cited (PTO-892)                                | 4) <input type="checkbox"/> Interview Summary (PTO-413)<br>Paper No(s)/Mail Date. _____ |
| 2) <input type="checkbox"/> Notice of Draftsperson's Patent Drawing Review (PTO-948)                       | 5) <input type="checkbox"/> Notice of Informal Patent Application                       |
| 3) <input type="checkbox"/> Information Disclosure Statement(s) (PTO/SB/08)<br>Paper No(s)/Mail Date _____ | 6) <input type="checkbox"/> Other: _____  |

### **DETAILED ACTION**

1. This action is in response to the papers filed September 10, 2007. Currently, claims 1-5, 12-14, 23-28, 31, 33-34 are pending.
2. All arguments have been thoroughly reviewed but are deemed non-persuasive for the reasons which follow. This action is made FINAL.
3. Any objections and rejections not reiterated below are hereby withdrawn.
  - a. The 103 rejections have been withdrawn in view of each of the claims to require detecting methylation of DAPK as characterizing or diagnosing breast cancer. The prior art does not teach an association between DAPK methylation and breast cancer.

### ***Priority***

4. This application claims priority to provisional 60/415,628, filed October 2, 2002.

### ***Drawings***

5. The drawings are acceptable.

### ***Claim Objections***

6. Claim 34 is objected to under 37 CFR 1.75(c), as being of improper dependent form for failing to further limit the subject matter of a previous claim. Applicant is required to cancel the claim(s), or amend the claim(s) to place the claim(s) in proper

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dependent form, or rewrite the claim(s) in independent form. Claim 34 depends on itself. Claim 34 depends on Claim 34 and has thus not been further treated on the merits.

**New Grounds of Rejection Necessitated by Amendments to the Claims**

***New Matter***

7. Claims 4, 24 are rejected under 35 U.S.C. 112, first paragraph, as containing subject matter which was not described in the specification in such a way as to reasonably convey to one skilled in the relevant art that the inventor(s), at the time the application was filed, had possession of the claimed invention.

In the amended claims, reference to “THBS” gene is included. The amendment proposes that the new claim language is supported by Figures 2 and 3. However, the specification does not describe or discuss “THBS” gene. Instead the specification describes a list of genes but fails to include THBS within this list (Figure 2, 3). This description does not support THBS gene. The concept of “THBS” gene does not appear to be part of the originally filed invention. Therefore, “THBS” gene constitutes new matter. Applicant is required to cancel the new matter in the reply to this Office Action.

***Claim Rejections - 35 USC § 112-Scope of Enablement***

The following is a quotation of the first paragraph of 35 U.S.C. 112:

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The specification shall contain a written description of the invention, and of the manner and process of making and using it, in such full, clear, concise, and exact terms as to enable any person skilled in the art to which it pertains, or with which it is most nearly connected, to make and use the same and shall set forth the best mode contemplated by the inventor of carrying out his invention.

8. Newly amended Claims 1-5, 12-14, 23-28, 31, 33-34 are rejected under 35

U.S.C. 112, first paragraph, because the specification, while being enabling for detecting methylation in DAPK genes, does not reasonably provide enablement for characterizing breast cancer based upon methylation profiles for DAPK. The specification does not enable any person skilled in the art to which it pertains, or with which it is most nearly connected, to use the invention commensurate in scope with these claims.

Factors to be considered in determining whether a disclosure meets the enablement requirement of 35 USC 112, first paragraph, have been described by the court in *In re Wands*, 8 USPQ2d 1400 (CA FC 1988). *Wands* states at page 1404,

“Factors to be considered in determining whether a disclosure would require undue experimentation have been summarized by the board in *Ex parte Forman*. They include (1) the quantity of experimentation necessary, (2) the amount of direction or guidance presented, (3) the presence or absence of working examples, (4) the nature of the invention, (5) the state of the prior art, (6) the relative skill of those in the art, (7) the predictability or unpredictability of the art, and (8) the breadth of the claims.”

The nature of the invention and breadth of claims

Claims 1-2, 23-28, 31 are drawn to a method of diagnosing breast cancer in a subject using plasma sample and detecting the presence or absence of DNA methylation in the promoters of a plurality of genes including DAPK.

Claims 3-5, 12-14 are drawn to method of characterizing cancer by providing a plasma sample from a subject and detecting the presence or absence of DNA

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methylation in DAPK to characterize breast cancer. Claim 5 illustrates characterize specifically encompasses diagnosing breast cancer.

Claim 33 is directed to diagnosing breast cancer in a subject using a biological sample and detecting methylation of DAPK to diagnose breast cancer.

The invention is in a class of invention which the CAFC has characterized as “the unpredictable arts such as chemistry and biology.” *Mycogen Plant Sci., Inc. v. Monsanto Co.*, 243 F.3d 1316, 1330 (Fed. Cir. 2001).

The unpredictability of the art and the state of the prior art

Maat et al. (Investigative Ophthalmology, Visual Science, Vol 48, No. 2, pages 486-490, February 2007) teaches a positive correlation was found between RASSF1a promoter methylation and development of metastatic disease, however a correlation with disease-free survival could not be established (abstract). Furthermore, Maat teaches that high frequency of RASSF1A methylation in cell lines compared with primary tumors was observed. Maat also teaches p16 methylation was more common in cell lines than tumors (page 489, col. 2). Thus Maat teaches that promoter methylation in genes is not indicative of any characterization and further that cell lines are not reliable predictors of tumor methylation.

Henrique et al. (Clin Cancer Research, Vol. 13, No. 20, pages 6122-6129, October 2007) teaches methylation patterns of different genes are associated with different characterizing events. For example, APC methylation appears to be associated with clinical Gleason score, however, CCND2, RARB2 are not associated with Gleason score. Henrique further teaches that APC hypermethylation was not significantly found to be associated with disease specific survival (page 6127, col. 1). Moreover, Henrique teaches that from a panel of five genes, CCND2 did not show a

significant association with disease-free survival in a univariate analysis (page 6128, col.2 ).

Suzuki et al. (Cancer Letters, Vol. 242, pages 222-230, 2006) teaches methylation patterns in cancer and finds that aberrant methylation differed between genes (see Table 4). Moreover, as seen in Figure 2, when analyzed in comparison to stages, different genes were found to have different patterns. Only one gene was significantly associated, namely CRBP1. Further, RIZ1 showed an inverse relationship from all the other genes. Thus it is unpredictable, without experimentation that is unpredictable whether genes are associated with stages or GS, for example.

Chang et al. (J. Mol. Med, Vol. 83, pages 132-139, 2005) teaches tamoxifen-resistant breast cancers show less frequent methylation of the estrogen receptor B but not the estrogen receptor alpha gene. Chang teaches that the methylation of ERB but not ERB was associated (abstract and Table 2). Thus, each gene is not similarly associated with resistance to drugs.

Maruya et al. (Clinical Cancer Research, Vol. 10, pages 3825-3830, June 2004) teaches cell lines and primary were analyzed and there was variability within and between cell lines and tumor specimens. This supports a heterogeneous and dynamic state of methylation in genes (abstract). Maruya clearly states that cell lines and carcinoma specimens manifest variable levels of methylation (page 3928, col. 2).

House (J. Gastrointest Surg, Vol. 7, pages 1004-1014, 2003) teaches analysis of promoter methylation in numerous genes. House teaches that statistical significance was reached only for tumor necrosis and E-cadherin gene methylation. Further, only E-cadherin methylation and absence of hMLH1 methylation correlated with early tumor recurrence. House analyzed the survival disadvantage at 5 years and found that the

promoter status of 10 tumor suppressor genes were not significant as prognostic markers (page 1009, col. 1-2).

The newly amended claims particularly claim plasma samples for detecting breast cancer. However, the post-filing date art makes it clear that DAPK is not specifically methylated in breast cancer. Yang et al (Gynecologic Oncology, Vol. 93, pages 435/-440, 2004) teaches detection of hypermethylated genes in tumor and plasma of cervical cancer patients. Table 2, page 437 illustrates, DAPK is methylated in plasma of cervical cancer patients.

Similarly, Miyamoto et al. (Jpn, J. Clin, Oncol, Vol. 35, No. 6, pages 293-301, 2005) teaches cancer-derived DNA in plasma by DNA methylation. DAPK and a plurality of other gene were analyzed in plasma and detected in Head and neck cancer (see Table 2, pages 296).

#### Guidance in the Specification.

The specification teaches methylation profiling in lymphoma cell lines (Figure 7). DAPK and PR are illustrated. For DAPK, 0% of the 8 controls have methylation while 33% of the T-cell lines were methylated. Similarly, for PR, 0% of the 8 control samples have methylation while 50% of the T-cell lines were methylated.

Table 1, page 76, illustrates differences between two cell lines. For example DAPK, was methylated in MCF7, but not methylated in T47D wt. Thus, cell lines do not appear to consistently demonstrate methylation.

The specification detects methylation in MDA-MD-231 breast cancer cell lines treated with 5-aza-2' deoxycytidine (page 77). The specification teaches analysis of 10 samples from patients. The specification fails to provide any results of the analysis, in particular with respect to DAPK or PR.



Figure 2 illustrates the methylation of 40 promoters in breast tumor tissue and normal breast tissue. As seen, the methylation of DAPK is unpredictable. DAPK is methylated in 5/5 normal tissues and in 4/5 breast tumor tissues (page 79). Moreover, PR is similarly methylated unpredictable.

Applicants have provided a declaration to support the teachings in the specification. Paragraph 4 of the declaration states that 29 plasma samples from normal patients or 29 samples from patients with DCIS were obtained. The declaration analyzes the methylation of DAPK1 in plasma of DCIS (ductal carcinomas in situ) patients and healthy controls. Table 3 of the declaration illustrates that DCIS is methylated 89.3% of the time whereas normal is methylated 51.9% of the time.

Table 4 of the declaration is described in paragraph 4 as teaching the biomarker comprising DAPK1 and additional genes was shown to identify DCIS with approximately 84% sensitivity and 90% specificity. The declaration fails to show which additional genes were used and how many additional genes, and whether any combination of genes would be significant.

Table 5 of the declaration is describes as plasma samples from three healthy patients or three plasma samples from patients with ADH (atypical ductal hyperplasia)(see paragraph 5 of the declaration). However the grant proposal appears to indicate that 8 samples of each were used. This appears to be a discrepancy between the explanation and the table. Moreover, ADH appears to be a benign condition and not be breast cancer, as particularly claimed in the pending claims. Finally the grant proposal indicates that "our experience indicates that genes with 20% methylation difference in the trial set almost always become components of the composite biomarker, suggesting that 18 genes can contribute to ADH biomarker." The

instant claims are not directed to a composite biomarker of 18 genes. Thus, the declaration does not appear to be commensurate in scope with the claims.

The guidance provided by the specification amounts to an invitation for the skilled artisan to try and follow the disclosed instructions to make and use the claimed invention.

#### Working Examples

The specification has no working examples of characterizing cancer for detecting chemotherapy resistant cancer, chance of disease free survival, risk of developing metastatic disease, monitoring progression.

#### Quantity of Experimentation

The quantity of experimentation in this area is extremely large since there is significant number of parameters which would have to be studied.

Claims 1, 3 and 5, for example are specifically drawn to diagnosing and characterizing breast cancer in a plasma sample by DAPK methylation patterns. The specification is silent with respect to any working examples of DAPK methylation patterns in plasma. The declaration filed April 29, 2008 appears to illustrate in DCIS patients, not any breast cancer, a difference between methylation in plasma of DCIS patients and healthy controls (89.3% v. 51.9%). It is noted there is no explicit p-value or statistical analysis for this biomarker. The analysis in the declaration for ADH has been considered, but does not appear to be within the scope of the claims, since the claims are drawn to breast cancer and not hyperplasia. Moreover, the sample size for the ADH analysis uses either 3 or 8 patients which is statistically small and lacks informative value. Finally, the claims appear to encompass a method of taking a plasma sample

from any subject, analyzing the plasma for methylation and assessing whether the individual has breast cancer. The post-filing date art suggest that at least two other cancers have high levels of methylation of DAPK in plasma. Thus, a method of screening patients for DAPK methylation in plasma would not indicate whether the subject had breast, head and neck or cervical cancer without further experimentation. Moreover, the absence of DNA methylation would not indicate an absence of breast cancer predictably. Breast cancer is affected by many genetic pathways which would similarly include BRAC1 or BRAC2 mutations. Thus, the subject may have an absence of methylation of DAPK but possess BRAC1 mutations and thus would have breast cancer. The specification and the declaration do not support the absence of methylation is indicative of the absence of breast cancer in the subject.

As previously noted, characterizing breast cancer, of Claim 3 encompasses detecting the presence or absence of chemotherapy resistant cancer. The specification fails to provide any guidance of how to characterize the cancer in such a manner. The art clearly teaches that different genes are differentially associated with resistance. Specifically, Chang teaches tamoxifen-resistant breast cancers show less frequent methylation of the estrogen receptor B but not the estrogen receptor alpha gene. Chang teaches that the methylation of ERB but not the ERA was associated (abstract and Table 2). Thus, each gene is not similarly associated with resistance to drugs. It would be unpredictable whether DAPK and PR are associated with chemotherapy resistant cancer because the state of the art teaches that without unpredictable and undue experimentation, different genes methylation patterns vary with respect to the association with resistance.

Similarly, characterizing breast cancer encompasses detecting a chance of disease-free survival, metastatic disease and progression. The specification fails to

provide any guidance of how to characterize the cancer in such a manner. The art clearly teaches that different genes are differentially associated with disease-free survival, metastatic disease and progression. The art teaches numerous situations where methylation patterns could not be established to be associated with a chance of disease-free survival, metastatic disease and progression. For example, Maat teaches a positive correlation was found between RASSF1a promoter methylation and development of metastatic disease, however a correlation with disease-free survival could not be established (abstract). Moreover, House teaches analysis of promoter methylation in numerous genes. House teaches that statistical significance was reached only for tumor necrosis and E-cadherin gene methylation. Further, only E-cadherin methylation and absence of hMLH1 methylation correlated with early tumor recurrence. House analyzed the survival disadvantage at 5 years and found that the promoter status of 10 tumor suppressor genes were not significant as prognostic markers (page 1009, col. 1-2). Therefore, for each particular gene and each particular “characterization” of disease-free survival, metastatic disease and progression, individual experimentation which is unpredictable and undue would be required. The experimentation would be trial and error experimentation with no expectation of success.

With respect to Claim 33 which is drawn to diagnosing breast cancer using any biological sample and DAPK methylation, the specification and the art clearly illustrate the unpredictability of cell lines and the differences in methylation. It is unpredictable given the small sample size with lack of significance whether any association between DAPK would exist without further experimentation. Maat teaches that high frequency of RASSF1A methylation in cell lines compared with primary tumors was observed. Maat also teaches p16 methylation was more common in cell lines than tumors (page 489,

col. 2). Thus Maat teaches that promoter methylation in cell lines are not reliable predictors of tumor methylation. Similarly, Maruya teaches cell lines and primary were analyzed and there was variability within and between cell lines and tumor specimens. This supports a heterogeneous and dynamic state of methylation in genes (abstract). Maruya clearly states that cell lines and carcinoma specimens manifest variable levels of methylation (page 3928, col. 2). Thus, analysis of cell lines without further experimentation in tumor cell lines would require unpredictable analysis regarding association of methylation patterns in tumor cells.

This would require significant inventive effort, with each of the many intervening steps, upon effective reduction to practice, not providing any guarantee of success in the succeeding steps.

#### Level of Skill in the Art

The level of skill in the art is deemed to be high.

#### Conclusion

In the instant case, as discussed above, in a highly unpredictable art where the art teaches the lack of association between genes and different characterizations including disease-free survival, metastatic disease and progression, the broad scope of the claims would require significant unpredictable and undue experimentation. Further, the prior art and the specification provides insufficient guidance to overcome the art recognized difficulties. Thus given the broad claims in an art whose nature is identified as unpredictable, the unpredictability of that art, the large quantity of research required to define these unpredictable variables, the lack of guidance provided in the

specification, the absence of a working example and the negative teachings in the prior art balanced only against the high skill level in the art, it is the position of the examiner that it would require undue experimentation for one of skill in the art to perform the method of the claim as broadly written.

### **Response to Arguments**

The response traverses the rejection. The response asserts the skilled artisan would be able to diagnose or characterized breast cancer in a subject as supported by the Declaration by inventor Dr. Vicro V. Levenson. The response asserts that the DAPK1 gene and additional genes were used together as a composite "biomarker". This argument has been considered but is not convincing because the declaration and specification do not provide the composite biomarkers which allow detection of breast cancer in plasma. The MPEP requires in 2164.05, "To overcome a prima facie case of lack of enablement, applicant must demonstrate by argument and/or evidence that the disclosure, as filed, would have enabled the claimed invention for one skilled in the art at the time of filing. This does not preclude applicant from providing a declaration after the filing date which demonstrates that the claimed invention works. However, the examiner should carefully compare the steps, materials, and conditions used in the experiments of the declaration with those disclosed in the application to make sure that they are commensurate in scope; i.e., that the experiments used the guidance in the specification as filed and what was well known to one of skill in the art. Such a showing also must be commensurate with the scope of the claimed invention, i.e., must bear a reasonable correlation to the scope of the claimed invention." Here, the declaration

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regarding the composite biomarker is not persuasive because the declaration and the specification do not provide such a composite biomarker and furthermore, the claims do not appear to be reasonably correlated to the scope of the claimed invention.

Thus for the reasons above and those already of record, the rejection is maintained.

### ***Claim Rejections - 35 USC § 112***

The following is a quotation of the second paragraph of 35 U.S.C. 112:

The specification shall conclude with one or more claims particularly pointing out and distinctly claiming the subject matter which the applicant regards as his invention.

9. Claims 35, 12-14 are rejected under 35 U.S.C. 112, second paragraph, as being indefinite for failing to particularly point out and distinctly claim the subject matter which applicant regards as the invention.

A) Claim 3 has been amended to require a plasma sample in step (a). However, the second line of step (a) remains drawn to "said biological sample". "Said biological sample lacks proper antecedent basis and thus is indefinite.

### ***Conclusion***

10. Applicant's amendment necessitated the new ground(s) of rejection presented in this Office action. Accordingly, **THIS ACTION IS MADE FINAL**. See MPEP § 706.07(a). Applicant is reminded of the extension of time policy as set forth in 37 CFR 1.136(a).

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A shortened statutory period for reply to this final action is set to expire THREE MONTHS from the mailing date of this action. In the event a first reply is filed within TWO MONTHS of the mailing date of this final action and the advisory action is not mailed until after the end of the THREE-MONTH shortened statutory period, then the shortened statutory period will expire on the date the advisory action is mailed, and any extension fee pursuant to 37 CFR 1.136(a) will be calculated from the mailing date of the advisory action. In no event, however, will the statutory period for reply expire later than SIX MONTHS from the date of this final action.

11. Any inquiry concerning this communication or earlier communications from the examiner should be directed to examiner Jeanine Goldberg whose telephone number is (571) 272-0743. The examiner can normally be reached Monday-Friday from 7:00 a.m. to 4:00 p.m.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Ram Shukla, can be reached on (571) 272-0735.

Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see <http://pair-direct.uspto.gov>. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free).

The Central Fax Number for official correspondence is (571) 273-8300.

/Jeanine A Goldberg/  
Primary Examiner, Art Unit 1634  
June 16, 2008